SAF-B00-054 100-NR-1 TSD Sites R. A. Sampling – Soil FINAL DATA PACKAGE

FAX R	ESULTS TO:		
	Rick Kerkow	372-8655	N/A INITIAL/DATE
VERIF	FICATION OF CL	IENT RECEI	PT:
	Phone or CC:Mail t	o Rick Kerkov	N/A
COMP	PLETE COPY OF	DATA PACK	AGE TO:
	Rick Kerkow	X5-60	BS 2/20/03 INITIALDATE
	Jeanette Duncan		BL SOV 103
COM! SHEE		E INCLUDE T	THE FOLLOWING ON THE FAX COVER
	SDG (H2051)	<u></u>	SAF-B00-054
	X Rad only	them only	Rad & Chem
	X Complete	Partial	

Waste Site: 116-N-1 Trench





February 15, 2003

Ms. Joan Kessner Bechtel Hanford Inc. 3350 George Washington Way Richland, WA 99352 MSIN: H0-25

Reference:

P.O. #630

Eberline Services R3-01-102-7756, SDG H2051

Dear Ms. Kessner:

Enclosed is the data report for three solid samples designated under SAF No. B00-054 received at Eberline Services on January 22, 2003. The samples were analyzed according to the accompanying chain-of-custody document.

Please call if you have any questions concerning this report.

Sincerely,

Melissa C. Mannion

Nelino Mamme

Program Manager

MCM

Enclosure: Data Package

FEE 2003

Case Narrative

Page 1 of 1

1.0 GENERAL

Bechtel Hanford Inc. (BHI) Sample Delivery Group H2051 was composed of three solid (soil) samples designated under SAF No. B00-054 with a Project Designation of: 100-NR-1 TSD Sites R.A. Sampling – Soil. The samples in SDG H2051 (Group R301102-7756) were batched with the samples in SDG H2050 (Group R301100-7436).

The samples were received as stated on the Chain-of-Custody documents. Any discrepancies are noted on the Eberline Services Sample Receipt Checklist. The results were transmitted to BHI via e-Fax on January 30, 2003.

2.0 ANALYSIS NOTES

2.1 Gross Alpha and Gross Beta Analyses

No problems were encountered during the course of the analyses.

2.2 Total Strontium Analyses

No problems were encountered during the course of the analyses.

2.3 Gamma Spectroscopy Analyses

No problems were encountered during the course of the analyses.

Case Narrative Certification Statement

"I certify that this data package is in compliance with the SOW, both technically and for completeness, for other than the conditions detailed above. Release of the data obtained in this hard copy data package has been authorized by the Laboratory Manager or a designee, as verified by the following signature."

Melissa C. Mannion
Program Manager

*A/15/3*Date

SDG <u>7756</u> Contact Melissa C. Mannion

Client <u>Hanford</u> Contract No. 630 Case no SDG H2051

SUMMARY DATA SECTION

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Melini Marm

Prepared by

Nelivi Mammi

Reviewed by

Lar id <u>EBRLNE</u> Protocol Hanford Version Ver 1.0 Form DVD-TOC Version 3.06 Report date <u>01/30/03</u>

SAMPLE DELIVERY GROUP H2051

SDG 7756
Contact Melissa C. Mannion

REPORT GUIDE

Client	Hanford
Contract	No. 630
Case no	SDG_H2051

ABOUT THE DATA SUMMARY SECTION

The Data Summary Section of a Data Package has all data, in several useful orders, necessary for first level, routine review of the data package for a Sample Delivery Group (SDG). This section follows the Data Package Narrative, which has an overview of the data package and a discussion of special problems. It is followed by the Raw Data Section, which has full details.

The Data Summary Section has several groups of reports:

SAMPLE SUMMARIES

The Sample and QC Summary Reports show all samples, including QC samples, reported in one SDG. These reports cross-reference client and lab sample identifiers.

PREPARATION BATCH SUMMARY

The Preparation Batch Summary Report shows all preparation batches (lab groupings reflecting how work was organized) relevant to the reported SDG with information necessary to check the completeness and consistency of the SDG.

WORK SUMMARY

The Work Summary Report shows all samples and work done on them relevant to the reported SDG.

METHOD BLANKS

The Method Blank Reports, one for each Method Blank relevant to the SDG, show all results and primary supporting information for the blanks.

LAB CONTROL SAMPLES

The Lab Control Sample Reports, one for each Lab Control Sample relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

REPORT GUIDES
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SUMMARY DATA SECTION

Page 1

SAMPLE DELIVERY GROUP H2051

SDG 7756
Contact Melissa C. Mannion

GUIDE, cont.

Client	Hani	ford	
Contract	No.	630	
Case no	SDG	H2051	

ABOUT THE DATA SUMMARY SECTION

DUPLICATES

The Duplicate Reports, one for each Duplicate and Original sample pair relevant to the SDG, show all results, differences and primary supporting information for these QC samples.

MATRIX SPIKES

The Matrix Spike Reports, one for each Spiked and Original sample pair relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

DATA SHEETS

The Data Sheet Reports, one for each client sample in the SDG, show all results and primary supporting information for these samples.

METHOD SUMMARIES

The Method Summary Reports, one for each test used in the SDG, show all results, QC and method performance data for one analyte on one or two pages. (A test is a short code for the method used to do certain work to the client's specification.)

REPORT GUIDES

The Report Guides, one for each of the above groups of reports, have documentation on how to read the associated reports.

REPORT GUIDES
Page 2
SUMMARY DATA SECTION
Page 2

SAMPLE DELIVERY GROUP H2051

SDG <u>7756</u> Contact Melissa C. Mannion

SAMPLE SUMMARY

Client <u>Hanford</u> Contract No. 630 Case no SDG H2051

CLIENT SAMPLE ID	LOCATION	MATRIX LEVEL	LAB SAMPLE ID	SAF NO	CHAIN OF CUSTODY	COLLECTED
J00F40	116-N-1 Trench	SOLID	R301102-01	B00-054	B00-054-225	01/16/03 08:50
J00F41	116-N-1 Trench	SOLID	R301102-02	B00-054	800-054-225	01/16/03 09:00
J00F42	116-N-1 Trench	SOLID	R301102-03	B00-054	B00-054-225	01/16/03 09:10
Method Blank		SOLID	R301100-03	B00-054		
Lab Control Sample		SOLID	R301100-02	B00-054		
Duplicate (R301102-01)	116-N-1 Trench	SOLID	R301102-04	B00-054		01/16/03 08:50

SAMPLE SUMMARY Page 1 SUMMARY DATA SECTION Page 3

SDG	7756	
Contact	<u>Melissa C.</u>	<u>Mannion</u>

QC SUMMARY

Client	Hanford	
Contract	No. 630	
Case no	SDG_H2051	

QC BATCH	CHAIN OF CUSTODY	CLIENT SAMPLE ID	MATRIX	% SOLIDS	SAMPLE AMOUNT	BAS1S AMOUNT	DAYS SI		LAB SAMPLE ID	DEPARTMENT SAMPLE ID
7436		Method Blank Lab Control Sample	SOL ID						R301100-03 R301100-02	7436-003 7436-002
7756	B00-054-225	J00F40	SOLID	88.9	1097 g		01/22/03	6	R301102-01	7756-001
		J00F41 J00F42	SOL ID	91.2 92.3	950 g 1093 g		01/22/03 01/22/03	6	R301102-02 R301102-03	7756-002 7756-003
		Duplicate (R301102-01)	SOLID	88.9	1097 g		01/22/03	6	R301102-04	7756-004

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Protocol Hanford
Version Ver 1.0
Form DVD-QS
Version 3.06
Report date 01/30/03

SDG	7756		
Contact	Melissa	С.	Mannion

PREP BATCH SUMMARY

Client	Hanford
Contract	No. 630
Case no	SDG H2051

TEST	MATRIX	METHOD	PREPARATION Batch		CLIENT	MORE		NCHETS Blank		DUP/ORIG MS/ORIG	QUALI- FIERS
Beta SR	Counting SOLID	Total Strontium in Soil	7043-061	10.0	3			1	1	1/1	
Gas F	roportion SOLID	al Counting Gross Alpha in Soil	7043-061	20.0	3	_		1	1	1/1	
93B	SOL ID	Gross Beta in Soil	7043-061	15.0	3		-	1	1	1/1	
GAM	Spectros SOLID	copy Gamma Scan	7043-061	15.0	3			1	1	1/1	

Duplicates and Matrix Spikes are those with original (Client) sample in this Sample Delivery Group.

Blank and LCS planchets are those in the same preparation batch as some Client, Duplicate or Spike sample.

PREP BATCH SUMMARY
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Protocol Hanford
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Version 3.06
Report date 01/30/03

SDG <u>7756</u> Contact <u>Melissa C. Mannion</u>

WORK SUMMARY

Client <u>Hanford</u>
Contract <u>No. 630</u>
Case no <u>SDG H2051</u>

CLIENT SAMPLE ID LOCATION CUSTODY	SAF No	MATRIX	LAB SAMPLE ID COLLECTED RECEIVED	PLANCHET	TEST	SUF-	ANALYZED	REVIEWED	ву	METHOD
J00F40			R301102-01	7756-001	93A/93		01/27/03	01/30/03	MCM	Gross Alpha in Soil
116-N-1 Trench		SOLID	01/16/03	7756-001	93B/93		01/27/03	01/30/03	MCM	Gross Beta in Soil
800-054-225	B00-054		01/22/03	7756-001	GAM		01/23/03	01/30/03	MCM	Gamma Scan
				7756-001	SR		01/27/03	01/30/03	MCM	Total Strontium in Soil
100F41			R301102-02	7756-002	93A/93		01/27/03	01/30/03	MCM	Gross Alpha in Soil
116-N-1 Trench		SOLID	01/16/03	7756-002	93B/93		01/27/03	01/30/03	MCM	Gross Beta in Soil
300-054-225	B00-054		01/22/03	7756-002	GAM		01/23/03	01/30/03	MCM	Gamma Scan
				7756-002	SR		01/27/03	01/30/03	MCM	Total Strontium in Soil
J00F42			R301102-03	7756-003	93A/93		01/27/03	01/30/03	MCM	Gross Alpha in Soil
116-N-1 Trench		SOLID	01/16/03	7756-003	938/93		01/27/03	01/30/03	MCM	Gross Beta in Soil
B00-054 - 225	B00-054		01/22/03	7756-003	GAM		01/23/03	01/30/03	MCM	Gamma Scan
				7756-003	SR		01/27/03	01/30/03	MCM	Total Strontium in Soil
lethod Blank			R301100-03	7436-003	93A/93	•	01/27/03	01/30/03	MCM	Gross Alpha in Soil
		SOLID		7436-003	93B/93		01/27/03	01/30/03	MCM	Gross Beta in Soil
	B00-054			7436-003	GAM		01/28/03	01/30/03	MCM	Gamma Scan
				7436-003	SR		01/27/03	01/30/03	MCM	Total Strontium in Soil
ab Control Sam	ple		R301100-02	7436-002	93A/93		01/28/03	01/30/03	MCM	Gross Alpha in Soil
		SOLID		7436-002	938/93		01/27/03	01/30/03	MCM	Gross Beta in Soil
	B00-054			7436-002	GAM		01/28/03	01/30/03	MCM	Gamma Scan
				7436-002	SR		01/27/03	01/30/03	MCM	Total Strontium in Soil
ouplicate (R301	102-01)		R301102-04	7756-004	93A/93		01/27/03	01/30/03	MCM	Gross Alpha in Soil
116-N-1 Trench		SOLID	01/16/03	7756-004	93B/93		01/27/03	01/30/03	MCM	Gross Beta in Soil
	B00-054		01/22/03	7756-004	GAM		01/24/03	01/30/03	MCM	Gamma Scan
				7756-004	SR		01/27/03	01/30/03	MCM	Total Strontium in Soil

WORK SUMMARY
Page 1
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SAMPLE DELIVERY GROUP H2051

SDG <u>7756</u> Contact <u>Melissa C. Mannion</u>

WORK SUMMARY, cont.

Client <u>Hanford</u>
Contract <u>No. 630</u>
Case no <u>SDG H2051</u>

TEST	SAF No	COUNTS	OF TESTS BY SA	CLIENT MORE	RE BLANK	LCS	DUP SPIKE	TOTAL
93A/93	B00-054	Gross Alpha in Soil	900.0_ALPHABETA_GPC	3	1	1	1	6
93B/93	B00-054	Gross Beta in Soil	900.0_ALPHABETA_GPC	3	1	1	1	6
GAM	800-054	Gamma Scan	GAMMA_GS	3	1	1	1	6
SR	B00-054	Total Strontium in Soil	SRTOT_SEP_PRECIP_GPC	3	1	1	1	6
TOTALS				12	4	4	4	24

WORK SUMMARY
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Form DVD-CWS
Version 3.06
Report date 01/30/03

R301100-03

METHOD BLANK

Method Blank

	7756 Melissa C. Mannion	Client/Case no Contract	 SDG_H2051
Lab sample id Dept sample id		Client sample id Material/Matrix SAF No	SOLID

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Gross Alpha	12587-46-1	-0.685	1.2	2.8	10	U	93A
Gross Beta	12587-47-2	0.023	3.3	5.6	15	ט	93B
Total Strontium	SR-RAD	0.075	0.47	0.97	1.0	U	SR
Potassium 40	13966-00-2	U		0.21		ט	GAM
Cobalt 60	10198-40-0	บ		0.011	0.050	U	GAM
Cesium 137	10045-97-3	ប		0.011	0.10	U	GAM
Radium 226	13982-63-3	U		0.019		υ	GAM
Radium 228	15262-20-1	U		0.048		U	GAM
Europium 152	14683-23-9	U		0.026	0.10	U	GAM
Europium 154	15585-10-1	U		0.029	0.10	U	GAM
Europium 155	14391-16-3	ΰ		0.030	0.10	U	GAM
Thorium 228	14274-82-9	U		0.014		ប	GAM
Thorium 232	TH-232	U		0.048		U	GAM
Uranium 235	15117-96-1	U		0.042		U	GAM
Uranium 238	U-238	U		1.4		บ	GAM
Americium 241	14596-10-2	บ		0.060		ט	GAM

100-NR-1 TSD Sites R.A. Sampling

QC-BLANK 43675

METHOD BLANKS
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R301100-02

LAB CONTROL SAMPLE

Lab Control Sample

SDG <u>7756</u> Contact <u>Melissa C. Mannion</u>	Client/Case no <u>Hanford SDG H2051</u> Contract <u>No. 630</u>
Lab sample id <u>R301100-02</u> Dept sample id <u>7436-002</u>	Client sample id <u>Lab Control Sample</u> Material/Matrix

ANALYTE	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCî/g	QUALI- FIERS	TEST	ADDED pCi/g	2σ ERR pCi/g	REC %	3σ LMTS (TOTAL)	PROTOCOL LIMITS
Gross Alpha	196	14	3.0	10		93A	200	8.0	98	68-132	70-130
Gross Beta	226	11	5.1	15		93B	212	8.5	107	74-126	70-130
Total Strontium	51.2	2.7	0.94	1.0		SR	53.0	2.1	97	83-117	80-120
Cobalt 60	0.280	0.019	0.010	0.050		GAM	0.299	0.012	94	76-124	80-120
Cesium 137	0.293	0.018	0.012	0.10		GAM	0.303	0.012	97	76-124	80-120

100-NR-1 TSD Sites R.A. Sampling

QC-LCS 43674		

LAB CONTROL SAMPLES
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R301102-04

DUPLICATE

J00F40

1	7756 Melissa C. Mannion		Client/Case no Contract	
İ	DUPLICATE	ORIGINAL		
Lab sample id	R301102-04	Lab sample id <u>R301102-01</u>	Client sample id	J00F40
Dept sample id	7756-004	Dept sample id <u>7756-001</u>	Location/Matrix	116-N-1 Trench SOLID
		Received <u>01/22/03</u>	Collected/Weight	01/16/03 08:50 1097 g
% solids	88.9	% solids <u>88.9</u>	Custody/SAF No	B00-054-225 B00-054

ANALYTE	DUPLICATE pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST	ORIGINAL pCi/g	2σ ERR (COUNT)	MDA pCi/g	QUALI- FIERS	RPD %	3σ τοτ	PROT Limit
Gross Alpha	0.714	1.9	3.1	10	U	93A	2.86	2.4	2.9	U	-		
Gross Beta	3.15	4.0	6.5	15	U	93B	1.56	3.3	5.5	U	-		
Total Strontium	0.029	0.12	0.25	1.0	U	SR	0.147	0.15	0.28	U	-		
Potassium 40	10.5	0.52	0.22			GAM	9.69	0.28	0.13		8	33	
Cobalt 60	U		0.027	0.050	υ	GAM	U		0.015	Ų	-		
Cesium 137	0.038	0.021	0.026	0.10		GAM	0.043	0.012	0.015		12	95	
Radium 226	0.367	0.048	0.046			GAM	0.334	0.026	0.024		9	40	
Radium 228	0.635	0.10	0.094			GAM	0.517	0.054	0.051		20	44	
Europium 152	U		0.058	0.10	U	GAM	u		0.031	U	_		
Europium 154	U		0.087	0.10	U	GAM	U		0.044	υ	-		
Europium 155	U		0.070	0.10	υ	GAM	υ		0.039	U	_		
Thorium 228	0.479	0.026	0.026			GAM	0.480	0.016	0.015		0	33	
Thorium 232	0.635	0.10	0.094			GAM	0.517	0.054	0.051		20	44	
Uranium 235	U		0.10		U	GAM	U		0.066	U	-		
Uranium 238	u		2.8		U	GAM	U		1.6	U	-		
Americium 241	U		0.15		U	GAM	U		0.087	U	_		

100-NR-1 TSD Sites R.A. Sampling

DUPLICATES
Page 1
SUMMARY DATA SECTION
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R301102-01

DATA SHEET

J00F40

)	7756 Melissa C. Mannion	Client/Case no Contract		SDG_H2051
		Client sample id Location/Matrix Collected/Weight Custody/SAF No	116-N-1 Trench 01/16/03 08:50	SOLID 1097 q B00-054

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Gross Alpha	12587-46-1	2.86	2.4	2.9	10	U	93A
Gross Beta	12587-47-2	1.56	3.3	5.5	15	Ū	93B
Total Strontium	SR-RAD	0.147	0.15	0.28	1.0	U	SR
Potassium 40	13966-00-2	9.69	0.28	0.13			GAM
Cobalt 60	10198-40-0	ប		0.015	0.050	ប	GAM
Cesium 137	10045-97-3	0.043	0.012	0.015	0.10		GAM
Radium 226	13982-63-3	0.334	0.026	0.024			GAM
Radium 228	15262-20-1	0.517	0.054	0.051			GAM
Europium 152	14683-23-9	Ū		0.031	0.10	U	GAM
Europium 154	15585-10-1	ΰ		0.044	0.10	U	GAM
Europium 155	14391-16-3	ŭ		0.039	0.10	บ	GAM
Thorium 228	14274-82-9	0.480	0.016	0.015			GAM
Thorium 232	TH-232	0.517	0.054	0.051			GAM
Uranium 235	15117-96-1	ប		0.066		บ	GAM
Uranium 238	U-238	Ŭ		1.6		υ	GAM
Americium 241	14596-10-2	Ū		0.087		Ū	GAM

100-NR-1 TSD Sites R.A. Sampling

DATA SHEETS
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SUMMARY DATA SECTION
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R301102-02

DATA SHEET

J00F41

	7756 Melissa C. Mannion	Client/Case no Contract	Hanford No. 630	SDG_	H2051
Lab sample id Dept sample id	•	Client sample id Location/Matrix			SOLID
Received	01/22/03 91.2	Collected/Weight Custody/SAF No	01/16/03 09:00	950 q B00-054	

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Gross Alpha	12587-46-1	1.92	2.4	3.3	10	ŭ	93A
Gross Beta	12587-47-2	-0.977	3.9	6.8	15	บ	93B
Total Strontium	SR-RAD	0.019	0.12	0.25	1.0	บ	SR
Potassium 40	13966-00-2	10.4	0.62	0.11			GAM
Cobalt 60	10198-40-0	0.034	0.013	0.014	0.050		GAM
Cesium 137	10045-97-3	0.080	0.009	0.010	0.10		GAM
Radium 226	13982-63-3	0.354	0.030	0.022			GAM
Radium 228	15262-20-1	0.574	0.059	0.052			GAM
Europium 152	14683-23-9	U		0.026	0.10	ט	GAM
Europium 154	15585-10-1	บ		0.036	0.10	U	GAM
Europium 155	14391-16-3	U		0.038	0.10	ប	GAM
Thorium 228	14274-82-9	0.518	0.017	0.012			GAM
Thorium 232	TH-232	0.574	0.059	0.052			GAM
Uranium 235	15117-96-1	U		0.053		U	GAM
Uranium 238	U-238	ប		1.8		U	GAM
Americium 241	14596-10-2	บ		0.015		υ	GAM

100-NR-1 TSD Sites R.A. Sampling

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R301102-03

DATA SHEET

J00F42

	7756 Melissa C. Mannion	Client/Case no Contract		DG_H2051
		Client sample id Location/Matrix Collected/Weight Custody/SAF No	116-N-1 Trench 01/16/03 09:10 1093	

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Gross Alpha	12587-46-1	2.54	2.5	3.4	10	υ	93A
Gross Beta	12587-47-2	2.40	3.6	5.9	15	U	93B
Total Strontium	SR-RAD	0.049	0.15	0.29	1.0	U	SR
Potassium 40	13966-00-2	9.56	0.37	0.20			GAM
Cobalt 60	10198-40-0	U		0.034	0.050	U	GAM
Cesium 137	10045-97-3	0.109	0.020	0.022	0.10		GAM
Radium 226	13982-63-3	0.360	0.032	0.032			GAM
Radium 228	15262-20-1	0.521	0.081	0.086			GAM
Europium 152	14683-23-9	บ		0.041	0.10	υ	GAM
Europium 154	15585-10-1	U		0.058	0.10	U	GAM
Europium 155	14391-16-3	บ		0.043	0.10	U	GAM
Thorium 228	14274-82-9	0.496	0.020	0.019			GAM
Thorium 232	TH-232	0.521	0.081	0.086			GAM
Uranium 235	15117-96-1	U		0.066		U	GAM
Uranium 238	U-238	ប		2.1		υ	GAM
Americium 241	14596-10-2	υ		0.058		Ŭ	GAM

100-NR-1 TSD Sites R.A. Sampling

DATA SHEETS
Page 3
SUMMARY DATA SECTION
Page 13

SAMPLE DELIVERY GROUP H2051

Test <u>SR</u> Matrix <u>SOLID</u>
SDG <u>7756</u>
Contact <u>Melissa C. Mannion</u>

METHOD SUMMARY

TOTAL STRONTIUM IN SOIL
BETA COUNTING

Client <u>Hanford</u>
Contract <u>No. 630</u>
Contract <u>SDG H2051</u>

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW SUF- TEST FIX	PLANCHET	Tot: Stron		
Preparation batch 7043-0	061	·				
J00F40	R301102-01		7756-001	U		
J00F41	R301102-02		7756-002	U		
J00F42	R301102-03		7756-003	U		
BLK (QC ID=43675)	R301100-03		7436-003	U		
LCS (QC 1D=43674)	R301100-02		7436-002	ok		
Duplicate (R301102-01)	R301102-04		7756-004	-	U	

METHOD PERFORMANCE

	LAB	RAW	SUF-	MDA	ALIQ	PREP	DILU-	YIELD	EFF	COUNT	FWHM	DRIFT	DAYS		ANAL-	
CLIENT SAMPLE ID	SAMPLE ID	TEST	FIX	pCi/g	g	FAC	TION	*	*	min	keV	KeV	HELD	PREPARED	YZED	DETECTOR
Preparation batch 7043-	061 2σ μ	rep er	ror 1	0.0 %	Reference	Lab	Notebool	k 7043	pg.	061						
J00F40	R301102-01			0.28	1.00			86		100			11	01/27/03	01/27	GRB-217
J00F41	R301102-02	2		0.25	1.00			93		100			11	01/27/03	01/27	GRB-218
J00F42	R301102-03	\$		0.29	1.00			84		100			11	01/27/03	01/27	GRB-221
BLK (QC 1D=43675)	R301100-03	3		0.97	0.400			73		100				01/27/03	01/27	GRB-224
LCS (QC ID=43674)	R301100-02	2		0.94	0.400			81		66				01/27/03	01/27	GRB-219
Duplicate (R301102-01)	R301102-04	•		0.25	1.00			91		100			11	01/27/03	01/27	GRB-222
(QC ID=43681)																
Nominal values and limi	ts from metl	ıod		1.0	0.400			30-10	5	100			180			

PROCEDURES	REFERENCE	SRTOT_SEP_PRECIP_GPC
	CP-060	Soil Preparation, rev 4
	CP-071	Soil Dissolution, > 1.0g Aliquot, rev 2
	CP-502	Strontium in Solids, rev 6

AVERAGES ± 2 SD	MDA	0.50	ŧ,	0.71
FOR 6 SAMPLES	YIELD .	85	±.	14

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SAMPLE DELIVERY GROUP H2051

Test 93A Matrix SOLID
SDG 7756
Contact Melissa C. Mannion

METHOD SUMMARY GROSS ALPHA IN SOIL GAS PROPORTIONAL COUNTING

Client Hanford
Contract No. 630
Contract SDG H2051

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW S TEST F	UF- IX PLANCHET	Gross	Alpha
Preparation batch 7043~	061				
J00F40	R301102-01	93	7756-001	U	
J00F41	R301102-02	93	7756-002	U	
J00F42	R301102-03	93	7756-003	U	
BLK (QC ID=43675)	R301100-03	93	7436-003	U	
LCS (QC 1D=43674)	R301100-02	93	7436-002	ok	
Duplicate (R301102-01)	R301102-04	93	7756-004	-	U

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW S	SUF- MD/ FIX pCi/g		PREP FAC		RESID	EFF %		FWHM keV		PREPARED	ANAL- YZED	DETECTOR
			po//s	·						***	 """	FALFARED	1200	DETECTOR
Preparation batch 7043-	061 2 <i>σ</i> pr	ep erre	or 20.0 %	Reference	Lab I	Notebook	7043	pg.	061					
J00F40	R301102-01	93	2.9	0.100			21		100		11	01/26/03	01/27	GRB-109
J00F41	R301102-02	93	3.3	0.100			21		100		11	01/26/03	01/27	GRB-110
J00F42	R301102-03	93	3.4	0.100			44		100		11	01/26/03	01/27	GRB-111
BLK (QC ID=43675)	R301100-03	93	2.8	0.100			21		100			01/26/03	01/27	GRB-102
LCS (QC ID=43674)	R301100-02	93	3.0	0.100			22		100			01/26/03	01/28	GRB-116
Duplicate (R301102-01)	R301102-04	93	3.1	0.100			22		100		11	01/26/03	01/27	GRB-112
(QC ID=43681)														
Nominal values and limi	ts from metho	od	10	0.100			5-250	0	100		 180		·	

PROCEDURES	REFERENCE	900.0_ALPHABETA_GPC
	CP-060	Soil Preparation, rev 4
	CP-071	Soil Dissolution, > 1.0g Aliquot, rev 2
	CP-125	Gross Alpha and Beta in Dissolved Solids, rev 3
L .		

AVERAGES ± 2 SD MDA 3.1 ± 0.46
FOR 6 SAMPLES RESIDUE 25 ± 18

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Protocol Hanford

Version Ver 1.0

Form DVD-CMS

Version 3.06

Report date 01/30/03

SAMPLE DELIVERY GROUP H2051

Test 938 Matrix SOLID
SDG 7756
Contact Melissa C. Mannion

METHOD SUMMARY GROSS BETA IN SOIL GAS PROPORTIONAL COUNTING

Client <u>Hanford</u>
Contract <u>No. 630</u>
Contract <u>SDG_H2051</u>

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW SU		Gross	s Beta
Preparation batch 7043-	061				
J00F40	R301102-01	93	7756-001	IJ	
J00F41	R301102-02	93	7756-002	U	
J00F42	R301102-03	93	7756-003	U	
BLK (QC ID=43675)	R301100-03	93	7436-003	Ų	
LCS (QC ID=43674)	R301100-02	93	7436-002	ok	
Duplicate (R301102-01)	R301102-04	93	7756-004	-	Ü

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST			PREP FAC	DILU- Tion	RESID mg	EFF %		 		PREPARED	ANAL- YZED	DETECTOR
Preparation batch 7043-0	061 2 <i>σ</i> pr	ep err	or 15.0 %	Reference	Lab I	Notebool	7043	pg.	061					
J00F40	R301102-01	93	5.5	0.100			21		100		11	01/26/03	01/27	GRB-109
J00F41	R301102-02	93	6.8	0.100			21		100		11	01/26/03	01/27	GRB-110
J00F42	R301102-03	93	5.9	0.100			44		100		11	01/26/03	01/27	GRB-111
BLK (QC ID=43675)	R301100-03	93	5.6	0.100			21		100			01/26/03	01/27	GRB-102
LCS (QC ID=43674)	R301100-02	93	5.1	0.100			22		100			01/26/03	01/27	GRB-101
Duplicate (R301102-01) (QC 1D=43681)	R301102-04	93	6.5	0.100			22		100		11	01/26/03	01/27	GRB-112
Nominal values and limi	ts from metho	xd	15	0.100			5-25	0	100		180	. <u> </u>		

PROCEDURES	REFERENCE	900.0_ALPHABETA_GPC
	CP-060	Soil Preparation, rev 4
	CP-071	Soil Dissolution, > 1.0g Aliquot, rev 2
	CP-125	Gross Alpha and Beta in Dissolved Solids, rev 3

AVERAGES ± 2 SD	MDA _	5.9	ŧ	1.3
FOR 6 SAMPLES	RESIDUE	25	±	18

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Protocol <u>Hanford</u>
Version <u>Ver 1.0</u>
Form <u>DVD-CMS</u>
Version <u>3.06</u>
Report date <u>01/30/03</u>

Lab id EBRLNE

SAMPLE DELIVERY GROUP #2051

Test GAM Matrix SOLID
SDG 7756
Contact Melissa C. Mannion

METHOD SUMMARY GAMMA SCAN GAMMA SPECTROSCOPY

Client Hanford
Contract No. 630
Contract SDG H2051

RESULTS

CLIENT SAMPLE ID		RAW SUF- TEST FIX PLANCHET	Cobalt 60	Cesium 137	
Preparation batch 7043-	061				
J00F40	R301102-01	7756-001	u	0.043	
J00F41	R301102-02	7756-002	0.034	0.080	
J00F42	R301102-03	7756-003	U	0.109	
BLK (QC ID=43675)	R301100-03	7436-003	u	U	
LCS (QC ID=43674)	R301100-02	7436-002	ok	ok	
Duplicate (R301102-01)	R301102-04	7756-004	- u	ok	
Nominal values and limit 100-NR-1 TSD Sites R.A.		RDLs (pCi/g)	0.050	0.10	

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB Sample	ID	RAW TEST	 		 DILU- Tion	YIELD			 		PREPARED	ANAL - YZED	DETECTOR
Preparation batch 7043-	061 2	e pi		 	Reference	 				 	THE LO	- NEI PRED	1220	DETECTOR
J00F40	R301102		•	0.10		 		F-3-	687		7	01/23/03	01/23	MB,05,00
J00F41	R301102	- 02		0.08	7 728				687		7	01/23/03	01/23	
J00F42	R301102	2-03		0.14	715				687			01/23/03		
BLK (QC 1D=43675)	R301100	- 03		0.08	6 772				260			01/23/03	-	• •
LCS (QC 1D=43674)	R301100	- 02		0.01	772				260			01/23/03		
Duplicate (R301102-01) (QC ID=43681)	R301102	2-04		0.19	_ 782				217		8	01/23/03		
Nominal values and limi	ts from m	neth	od	 0.05	0 772	 			100	 	180			

PROCEDURES	REFERENCE	GAMMA_GS
	CP-060	Soil Preparation, rev 4
	CP-100	Ge(Li) Preparation for Commercial Samples, rev 5

AVERAGES ± 2 SD	MDA <u>0.10</u> ± <u>0.12</u>
FOR 6 SAMPLES	YIELD ±

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SAMPLE DELIVERY GROUP H2051

SDG 7756
Contact Melissa C. Mannion

REPORT GUIDE

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SAMPLE SUMMARY

The Sample and QC Summary Reports show all samples, including QC samples, reported in one Sample Delivery Group (SDG).

The Sample Summary Report fully identifies client samples and gives the corresponding lab sample identification. The QC Summary Report shows at the sample level how the lab organized the samples into batches and generated QC samples. The Preparation Batch and Method Summary Reports show this at the analysis level.

The following notes apply to these reports:

- * LAB SAMPLE ID is the lab's primary identification for a sample.
- * DEPARTMENT SAMPLE ID is an alternate lab id, for example one assigned by a radiochemistry department in a lab.
- * CLIENT SAMPLE ID is the client's primary identification for a sample. It includes any sample preparation done by the client that is necessary to identify the sample.
- * QC BATCH is a lab assigned code that groups samples to be processed and QCed together. These samples should have similar matrices.

QC BATCH is not necessarily the same as SDG, which reflects samples received and reported together.

* All Lab Control Samples, Method Blanks, Duplicates and Matrix Spikes are shown that QC any of the samples. Due to possible reanalyses, not all results for all these QC samples may be relevant to the SDG. The Lab Control Sample, Method Blank, Duplicate, Matrix Spike and Method Summary Reports detail these relationships.

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SAMPLE DELIVERY GROUP H2051

SDG 7756
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PREPARATION BATCH SUMMARY

The Preparation Batch Summary Report shows all preparation batches in one Sample Delivery Group (SDG) with information necessary to check the completeness and consistency of the SDG.

The following notes apply to this report:

- * The preparation batches are shown in the same order as the Method Summary Reports are printed.
- * Only analyses of planchets relevant to the SDG are included.
- * Each preparation batch should have at least one Method Blank and LCS in it to validate client sample results.
- * The QUALIFIERS shown are all qualifiers other than U, J, B, L and H that occur on any analysis in the preparation batch. The Method Summary Report has these qualifiers on a per sample basis.

These qualifiers should be reviewed as follows:

- X Some data has been manually entered or modified. Transcription errors are possible.
- P One or more results are 'preliminary'. The data is not ready for final reporting.
- 2 There were two or more results for one analyte on one planchet imported at one time. The results in DVD may not be the same as on the raw data sheets.

Other lab defined qualifiers may occur. In general, these should be addressed in the SDG narrative.

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SAMPLE DELIVERY GROUP H2051

SDG <u>7756</u>
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WORK SUMMARY

The Work Summary Report shows all samples, including QC samples, and all relevant analyses in one Sample Delivery Group (SDG). This report is often useful as supporting documentation for an invoice.

The following notes apply to this report:

- * TEST is a code for the method used to measure associated analytes. Results and related information for each analyte are on the Data Sheet Report. In special cases, a test code used in the summary data section is not the same as in associated raw data. In this case, both codes are shown on the Work Summary.
- * SUFFIX is the lab's code to distinguish multiple analyses (recounts, reworks, reanalyses) of a fraction of the sample. The suffix indicates which result is being reported. An empty suffix normally identifies the first attempt to analyze the sample.
- * The LAB SAMPLE ID, TEST and SUFFIX uniquely identify all supporting data for a result. The Method Summary Report for each TEST has method performance data, such as yield, for each lab sample id and suffix and procedures used in the method.
- * PLANCHET is an alternate lab identifier for work done for one test. It, combined with the TEST and SUFFIX, may be the best link to raw data.
- * For QC samples, only analyses that directly QC some regular sample are shown. The Lab Control Sample, Method Blank, Duplicate, Matrix Spike and Method Summary Reports detail these relationships.
- * The SAS (Special Analytical Services) Number is a client or lab assigned code that reflects special processing for samples, such as rapid turn around. Counts of tests done are lists by SAS number since it is likely to affect prices.

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DATA SHEET

The Data Sheet Report shows all results and primary supporting information for one client sample or Method Blank. This report corresponds to both the CLP Inorganics and Organics Data Sheet.

The following notes apply to this report:

- * TEST is a code for the method used to measure an analyte. If the TEST is empty, no data is available; the analyte was not analyzed for.
- * The LAB SAMPLE ID and TEST uniquely identify work within the Summary Data Section of a Data Package. The Work Summary and Method Summary Reports further identify raw data that underlies this work.

The Method Summary Report for each TEST has method performance data, such as yield, for each Lab Sample ID and a list of procedures used in the method.

- * ERRORs can be labeled TOTAL or COUNT. TOTAL implies a preparation (non-counting method) error has been added, as square root of sum of squares, to the counting error denoted by COUNT. The preparation errors, which may vary by preparation batch, are shown on the Method Summary Report.
- * A RESULT can be 'N.R.' (Not Reported). This means the lab did this work but chooses not to report it now, possibly because it was reported at another time.
- * When reporting a Method Blank, a RESULT can be 'N.A.' (Not Applicable). This means there is no reported client sample work in the same preparation batch as the Blank's result. This is likely to occur when the Method Blank is associated with reanalyses of selected work for a few samples in the SDG.

The following qualifiers are defined by the DVD system:

U The RESULT is less than the MDA (Minimum Detectable Activity).

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DATA SHEET

If the MDA is blank, the ERROR is used as the limit.

- J The RESULT is less than the RDL (Required Detection Limit) and no U qualifier is assigned.
- B A Method Blank associated with this sample had a result without a U flag and, after correcting for possibly different aliquots, that result is greater than or equal to the MDA for this sample.

Normally, B is not assigned if U is. When method blank subtraction is shown on this report, B flags are assigned based on the unsubtracted values while U's are assigned based on the subtracted ones. Both flags can be assigned in this case.

For each sample result, all Method Blank results in the same preparation batch are compared. The Method Summary Report documents this and other QC relationships.

- L Some Lab Control Sample that QC's this sample had a low recovery. The lab can disable assignment of this qualifier.
- H Similar to 'L' except the recovery was high.
- P The RESULT is 'preliminary'.
- X Some data necessary to compute the RESULT, ERROR or MDA was manually entered or modified.
- There were two or more results available for this analyte. The reported result may not be the same as in the raw data.

Other qualifiers are lab defined. Definitions should be in the SDG narrative.

The following values are underlined to indicate possible problems:

* An MDA is underlined if it is bigger than its RDL.

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DATA SHEET

- * An ERROR is underlined if the 1.645 sigma counting error is bigger than both the MDA and the RESULT, implying that the MDA may not be a good estimate of the 'real' minimum detectable activity.
- * A negative RESULT is underlined if it is less than the negative of its 2 sigma counting ERROR.
- * When reporting a Method Blank, a RESULT is underlined if greater than its MDA. If the MDA is blank, the 2 sigma counting error is used in the comparison.

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REPORT GUIDE

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LAB CONTROL SAMPLE

The Lab Control Sample Report shows all results, recoveries and primary supporting information for one Lab Control Sample.

The following notes apply to this report:

- * All fields in common with the Data Sheet Report have similar usage. Refer to its Report Guide for details.
- * An amount ADDED is the lab's value for the actual amount spiked into this sample with its ERROR an estimate of the error of this amount.

An amount added is underlined if its ratio to the corresponding RDL is outside protocol specified limits.

- * REC (Recovery) is RESULT divided by ADDED expressed as a percent.
- * The first, computed limits for the recovery reflect:
 - 1. The error of RESULT, including that introduced by rounding the result prior to printing.

If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.

- 2. The error of ADDED.
- 3. A lab specified, per analyte bias. The bias changes the center of the computed limits.
- * The second limits are protocol defined upper and lower QC limits for the recovery.
- * The recovery is underlined if it is outside either of these ranges.

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DUPLICATE

The Duplicate Report shows all results, differences and primary supporting information for one Duplicate and associated Original sample.

The following notes apply to this report:

* All fields in common with the Data Sheet Report have similar usage. This applies both to the Duplicate and Original sample data. Refer to the Data Sheet Report Guide for details.

If the Duplicate has data for a TEST and the lab did not do this test to the Original, the Original's RESULTs are underlined.

* The RPD (Relative Percent Difference) is the absolute value of the difference of the RESULTs divided by their average expressed as a percent.

If both RESULTs are less than their MDAs, no RPD is computed and a '-' is printed.

For an analyte, if the lab did work for both samples but has data for only one, the MDA from the sample with data is used as the other's result in the RPD.

* The first, computed limit is the sum, as square root of sum of squares, of the errors of the results divided by the average result as a percent, hence the relative error of the difference rather than the error of the relative difference. The errors include those introduced by rounding the RESULTs prior to printing.

If this limit is labeled TOT, it includes the preparation error in the RESULTs. If labeled CNT, it does not.

This value reported for this limit is at most 999.

- * The second limit for the RPD is the larger of:
 - 1. A fixed percentage specified in the protocol.

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DUPLICATE

- A protocol factor (typically 2) times the average MDA as a percent of the average result. This limit applies when the results are close to the MDAs.
- * The RPD is underlined if it is greater than either limit.
- * If specified by the lab, the second limit column is replaced by the Difference Error Ratio (DER), which is the absolute value of the difference of the results divided by the quadratic sum of their one sigma errors, the same errors as used in the first limit.

Except for differences due to rounding, the DER is the same as the RPD divided by the first RPD limit with the limit scaled to 1 sigma.

* The DER is underlined if it is greater than the sigma factor, typically 2 or 3, shown in the header for the first RPD limit.

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SAMPLE DELIVERY GROUP H2051

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MATRIX SPIKE

The Matrix Spike Report shows all results, recoveries and primary supporting information for one Matrix Spike and associated Original sample.

The following notes apply to this report:

- * All fields in common with the Data Sheet Report have similar usage. This applies both to the Spiked and Original sample data. Refer to the Data Sheet Report Guide for details.
 - If the Spike has data for a TEST and the lab did not do this test to the Original, the Original's RESULTs are underlined.
- * An amount ADDED is the lab's value for the actual amount spiked into the Spike sample with its ERROR an estimate of the error of this amount.
 - An amount is underlined if its ratio to the corresponding RDL is outside protocol specified limits.
- * REC (Recovery) is the Spike RESULT minus the Original RESULT divided by ADDED expressed as a percent.
- * The first, computed limits for the recovery reflect:
 - 1. The errors of the two RESULTs, including those introduced by rounding them prior to printing.
 - If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.
 - 2. The error of ADDED.
 - 3. A lab specified, per analyte bias. The bias changes the center of the computed limits.
- * The second limits are protocol defined upper and lower QC limits

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SAMPLE DELIVERY GROUP H2051

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MATRIX SPIKE

for the recovery.

These limits are left blank if the Original RESULT is more than a protocol defined factor (typically 4) times ADDED. This is a way of accounting for that when the spike is small compared to the amount in the original sample, the recovery is unreliable.

* The recovery is underlined (out of spec) if it is outside either of these ranges.

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SAMPLE DELIVERY GROUP H2051

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METHOD SUMMARY

The Method Summary Report has two tables. One shows up to five results measured using one method. The other has performance data for the method. There is one report for each TEST, as used on the Data Sheet Report.

The following notes apply to this report:

* Each table is subdivided into sections, one for each preparation batch. A preparation batch is a group of aliquots prepared at roughly the same time in one work area of the lab using the same method.

There should be Lab Control Sample and Method Blank results in each preparation batch since this close correspondence makes the QC meaningful. Depending on lab policy, Duplicates need not occur in each batch since they QC sample dependencies such as matrix effects.

* The RAW TEST column shows the test code used in the raw data to identify a particular analysis if it is different than the test code in the header of the report. This occurs in special cases due to method specific details about how the lab labels work.

The Lab Sample or Planchet ID combined with the (Raw) Test Code and Suffix uniquely identify the raw data for each analysis.

* If a result is less than both its MDA and RDL, it is replaced by just 'U' on this report. If it is greater than or equal to the RDL but less than the MDA, the result is shown with a 'U' flag.

The J and X flags are as on the data sheet.

- * Non-U results for Method Blanks are underlined to indicate possible contamination of other samples in the preparation batch. The Method Blank Report has supporting data.
- * Lab Control Sample and Matrix Spike results are shown as: ok, No data, LOW or HIGH, with the last two underlined. 'No data'

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SAMPLE DELIVERY GROUP H2051

SDG 7756
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METHOD SUMMARY

means no amount ADDED was specified. 'LOW' and 'HIGH' correspond to when the recovery is underlined on the Lab Control Sample or Matrix Spike Report. See these reports for supporting data.

- * Duplicate sample results are shown as: ok, No data, or OUT, with the last two underlined. 'No data' means there was no original sample data found for this duplicate. 'OUT' corresponds to when the RPD is underlined on the Duplicate Report. See this report for supporting data.
- * If the MDA column is labeled 'MAX MDA', there was more than one result measured by the reported method and the MDA shown is the largest MDA. If not all these results have the same RDL, the MAX MDA reflects only those results with RDL equal to the smallest one.

MDAs are underlined if greater than the printed RDL.

- * Aliquots are underlined if less than the nominal value specified for the method.
- * Prepareation factors are underlined if greater than the nominal value specified for the method.
- * Dilution factors are underlined if greater than the nominal value specified for the method.
- * Residues are underlined if outside the range specified for the method. Residues are not printed if yields are.
- * Yields, which may be gravimetric, radiometric or some type of recovery depending on the method, are underlined if outside the range specified for the method.
- * Efficiencies are underlined if outside the range specified for the method. Efficiencies are detector and geometry dependent so this test is only approximate.

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SAMPLE DELIVERY GROUP H2051

SDG 7756
Contact Melissa C. Mannion

GUIDE, cont.

Client	Hanf	ord	
Contract	No.	630	
Case no	SDG	H2051	

METHOD SUMMARY

- * Count times are underlined if less than the nominal value specified for the method.
- * Resolutions (as FWHM; Full Width at Half Max) are underlined if greater than the method specified limit.
- * Tracer drifts are underlined if their absolute values are greater than the method specified limit. Tracer drifts are not printed if percent moistures are.
- * Days Held are underlined if greater than the holding time specified in the protocol.
- * Analysis dates are underlined if before their planchet's preparation date or, if a limit is specified, too far after it.

For some methods, ratios as percentages and error estimates for them are computed for pairs of results. A ratio column header like '1 \div 3' means the ratio of the first result column and the third result column.

Ratios are not computed for Lab Control Sample, Method Blank or Matrix Spike results since their matrices are not necessarily similar to client samples'.

The error estimate for a ratio of results from one planchet reflects only counting errors since other errors should be correlated. For a ratio involving different planchets, if QC limits are computed based on total errors, the error for the ratio allows for the preparation errors for the planchets.

The ratio is underlined (out of spec) if the absolute value of its difference from the nominal value is greater than its error estimate. If no nominal value is specified, this test is not done.

For Gross Alpha or Gross Beta results, there may be a column showing the sum of other Alpha or Beta emitters. This sum includes all relevant

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SAMPLE DELIVERY GROUP H2051

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GUIDE, cont.

Client	Hani	ord
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Case no	<u>\$DG</u>	H2051

METHOD SUMMARY

results in the DVD database, whether reported or not. Results in the sum are weighted by a particles/decay value specified by the lab for each relevant analyte. Results less than their MDA are not included. No sums are computed for Lab Control, Method Blank or Matrix Spike samples since their various planchets may not be physically related.

If a ratio of total isotopic to Gross Alpha or Beta is shown, the error for the ratio reflects both the error in the Gross result and the sum, as square root of sum of squares, of the errors in the isotopic results.

For total elemental uranium or thorium results, there may be a column showing the total weight computed from associated isotopic results. Ignoring results less than their MDAs, this is a weighted sum of the isotopic results. The weights depend on the molecular weight and half-life of each isotope so as to convert activities (decays) to weight (atoms).

If a ratio of total computed to measured elemental uranium or thorium is shown, the error for the ratio reflects the errors in all the measurements.

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JODF42 SOIL 1-16-03 091D X CHAIN OF POSSESSION Sign/Print Names Relinquished By/Remand From Box 1-16-03 Received Try/Stored In Posterior 1-16-03 Received Try/Stored In Posterior 1-16-03 Received By/Stored In Received By/Stored In Posterior 1-16-03 Delett: 15-07-091 Delet																_	
RR Kechow Treford Determination 100 NR: TSO Sities R. A Sampling - Soil Sampling Location 110 NR: TSO Sities R. A Sampling - Soil Siting of To The Christ Nr. Profest Determination 110 NR: TSO Sities R. A Sampling - Soil Siting of To The Christ Nr. Profest Determination 110 NR: TSO Sities R. A Sampling - Soil Siting of To The Christ Nr. Profest Determination Risk Treets Risk Tree	Bechtel Hanford Inc. CHAIN OF CUSTODY/SAMPI					PLE	LE ANALYSIS REQUEST				B00-054-225 P		Page 1	of 1			
Project Designation Sampling Scale Sampling Sam										dinator	Price Code	3]	Data Tu	rnaround			
Fee Chest No. -9						g Location	2051	((77	56)				_	-	7 days	
Sample No. Sample No. Sample No. JODE 40 SOIL JODE 40 SEPECIAL INSTRUCTIONS LAB COAL ISSUMMENT JOURN SEPECIAL INSTRUCTIONS J	Ice Chest No.	TJ-9	3	1		_			-	00		Method of Sh		E			
POSSIBLE SAMPLE HATARDSREMARKS Proteinly reductive SEE BELOW FOR DCTIVITY Report Type of Container No. of Container(s) No. of Container(s) No. of Container(s) Sample No. Sample No. Matrix * Sample Date Special Insurance Insu			<u></u>	Of	fsite F	Property No.	1030	12	0			Bill of Ladin		No.	8019	3 9/081	()
SEE BEIOU FOR ACTIVITY Report ### Type of Container Type of Con			S/REMARKS		T					·	<u> </u>		1			,_,_	Ī
SPECIAL INSTRUCTIONS Reliquished By-Removed From Date/Time Received By/Sored in Date/Time Re	Potentially radioa	ctive				Preservation	None								•		
SAMPLE ANALYSIS Sample No. Matrix * Sample Date Sample Time Special Information Social State of the Sample Time Special Information Social So	SEE REL	OW FOR 1	CTIVITY R	aport #	- -	Trest, value	Marinelli	Ne.			 			- 	}		
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Sample No. Matrix * Sample Date Sample Time Some land Interactions Sample No. Matrix * Sample Date Sample Time Social Interactions Solid Interactions Interactions Solid Interactions Solid Interactions Solid Interactions Solid Interactions Interactions Solid Interactions Interactions Solid Interactions Solid Interactions Solid Interactions Solid Interactions Solid Interactions Interactions Solid Interactions Solid Interactions Solid Interactions Solid Interactions Solid Interactions Solid Interactions Interactions Solid Interactions Interactions Interactions Interactions	1 -	•				No. of Container(s)	1			J]	}]	j]	j	
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JODF40 SOIL JODF41 SOIL JODF42 SOIL JODF42 SOIL JODF42 SOIL JODF43 SIgn/Print Names Relinquished By/Removed From Date/Time Received By/Stored In Date/Time Date/Time Received By/Stored In Date/Time Title Date/Time														THETO			
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JODF41 SOIL (-16-03 0900 X JODF49 63-0 JODF42 SOIL (-16-03 0910 X JODF30 GB-0 CHAIN OF POSSESSION Sign/Print Names Reliquished By/Removed From Date/Time Received By/Stored In Date/Time Reliquished By/Removed From Date/Time Received By/Stored In Date/Time	J00F40		SOIL	1-16-	0)	0350	X							100F28			6B-01
CHAIN OF POSSESSION Sign/Print Names Relinquished By/Removed From Pater I A 3728 Relinquished By/Removed From Pater I By/Stored In Received By/Stored In Received By/Stored In Pater I By/Stored In Received By/Stored In Pater I By/Sto	J00F41		SOIL	1-16-	03		X							J00129			613-02
Relinquished By/Removed From Date/Time Received By/Stored In Date/Time Date/Time Date/Time Date/Time Received By/Stored In Date/Time Date/Time Date/Time Date/Time Date/Time Received By/Stored In Date/Time Date/Ti	J00F42		SOIL											J00F30			6B-03
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Relinquished By/Removed From Date/Time Received By/Stored In Date/Time LAB COA: R1301N2F00	CHAIN OF P	POSSESSION						<u> </u>	SPEC	IAL INSTR	UCTIO	NS					Matrix *
Relinquished By/Removed From Pate/Time Date/Time 			Received By/	Stored	In ASher Do	1-16-03	<u> </u>	1			sium-137. Cobalt	-60. Europiu	m-152. Euronium-	154, Europium	-155};	SE=Sediment	
Relinquished By/Removed From Date/Time Received By/Stored In Date/Time Personnel not available to relinquish samples from the 3728 Ref # 1/2 on 1 / 21 / 2	Relinquished By/Remo	ved From		Received By/	Stored	in Da	ite/Time 100	<u>.</u>	Correct	a Spec - Adda	on LAmeri	rinm-241 lasenin	ric Physician	n American 241	: Strontium-89.	90 Total	Si=Sludge
Relinquished By/Removed From Date/Time Received By/Stored In Date/Time Personnel not available to relinquish samples from the 3728 Ref # 1/2 on 1 / 21 / 2	Relinquished By/Remov	ved From	Date/Time 1100	Received By			zi-o3 ue/Time		DE	ETE:	1507	OPIC PU	, Am	241 , Ni	63, 7	unum	
Relinquished By/Removed From Date/Time Received By/Sured In Date/Time relinquish samples from the 3728 Ref # 1/2 on 1 / 21 / 2				Received By/	Stored				n	annal not a	veitable i	io.		1-16-03		1	T=Tissue WI=Wipe L=Liquid
LABORATORY Received By SECTION Title Date/Time FINAL SAMPLE Disposal Method	Relinquished By/Remov	ved From	Date/Time						relii	rauish samp	oles from	the 3728	'				
LABORATORY SECTION Received By The Disposed By Date/Time	Relinquished By/Remov	ved From	Date/Time	Received By/	Stored	In Da	ite/Time				<u>.</u>						
FINAL SAMPLE Disposal Method Disposal Method Disposed By		Received By					Tit	le							Di	ate/Time	
	FINAL SAMPLE	Disposal Method								Dispos	sed By			,	D		

Richmond, CA Laboratory

SAMPLE RECEIPT CHECKLIST

		SAMPLE RECEI	PT		
Client:	BHI	Dat	e/Time receive	d 1-22-05	3/945:
CoC N	o. <u>B06 054-225</u>			•	<i>'</i> .
1)	ner I.D. No. TJ 9	Requested TAT (D	avs) 7 P.	O. Received	Yes! No.! 1
		INSPECTION			rea () NO ()
	Custody seals on shipping cor	•	Yes [(]	No[]	NIA F 3
1. 2.	Custody seals on shipping cor			No []	N/A []
3.	Custody seals on sample cont	•	Yes [/]	No []	N/A []
4.	Custody seals on sample conf		,	No []	N/A [·]
5.	Packing material is:	and dated a sign	Wet [/]	Dry []	N/A []
6.	Number of semples in shippin	a container:	_	D, y ()	·
7.	Number of containers per san	rple:	(Or see Co	nC	
8.	Paperwork agrees with sample		· ·	No []	- 1
9.	Samples have: Tape [] Ha				le labels f
10.	Samples are: in good condit				
11.		-			
1					
#					
13.	Was P.M. notified of any and	malies? Yes []	No []	Date	
14.	Received by www.	, Da	ate: 1-22-0	<u> ろ</u> Time: (2	141
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Custom	er Sample (No. cpm mr/hr	Custo wipe	omer Sample No.	cpm m	ır/hr wipe
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Survey N	Meter Ser. No.	Calibr	ation date		
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Form SCP-01.2, 05-03-02

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